From: Sent: To: Subject:

Pak, Yong Tuesday, January 15, 2002 1:38 PM STIC-Biotech/ChemLib 09/654,652

dear stic,

please search the following in commercial and interference database for 09/654,652:

- 1. SEQ ID NO:3
- 2. oligomer search of SEQ ID NO:3

Yong Pak

Art Unit 1652 CM1 10A16 (mail box 10C01) 703-308-9363

Pending Nucleic Acid and/or Pending Amino Acid database searches now generate two sets of results. These databases were split into to two parts to reduce the time needed to update the databases daily. The split freed up more machine time for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions, .rnpm and .rnpn

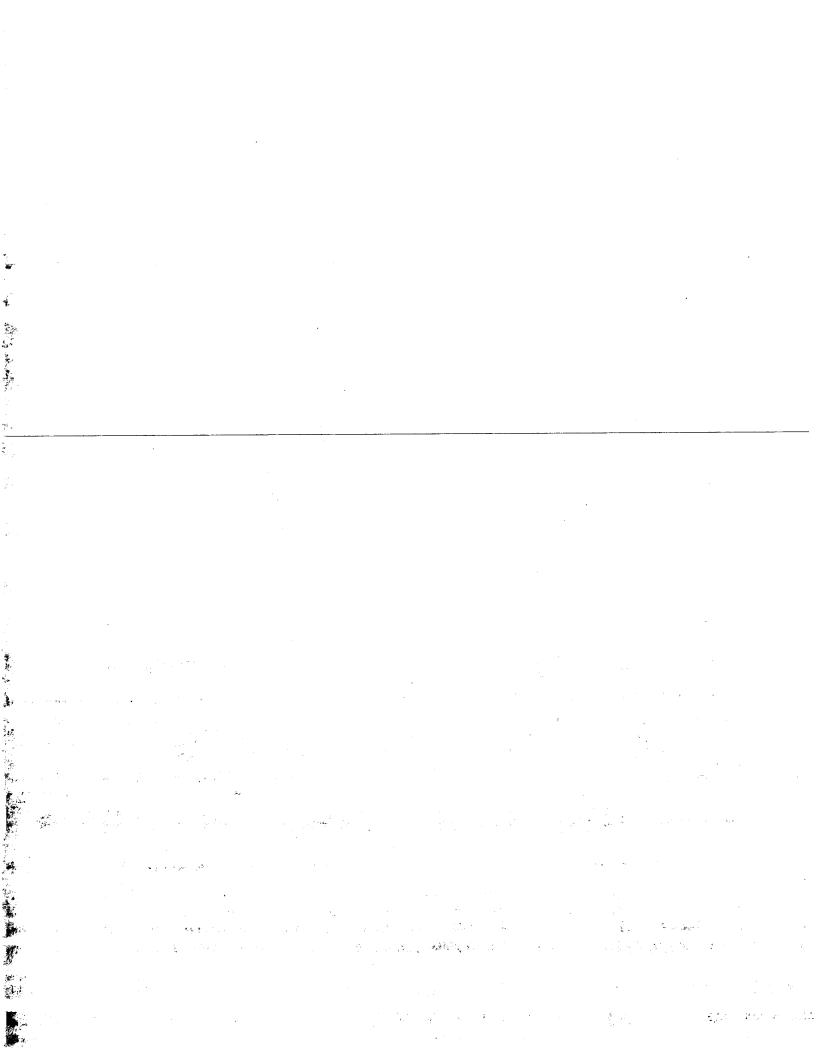
Searches run against the Amino Acid Pending database produce two sets of results, with the extensions, .rapm and .rapn

The Pending database search results should not be left in the case because they contain data that is confidential.

Searcher: Alessa- Vall
Phone: 308-450/
Location: Sinter lub.
Date Picked Up: 1/15/02
Date Completed: 1/15-102
Searcher Prep/Review:
Clerical: 57
Online time: 4

TYPE OF SEARCH:
NA Sequences:
AA Sequences: 2
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST(where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.: ABSSO2
WWW/Internet:
Other (specify):



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Title:
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EBARWKMAAASGTVS 60	ength 228; Indels 0; Gaps		Sequence 6, Appli Sequence 16, Appli Sequence 3, Appli Sequence 3, Appli Sequence 3, Appli Sequence 1, Appli Sequence 1, Appli Sequence 3, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 2, Appli

RESULT 2
US-08-103-998-2
; Sequence 2, Application US/08103998
; Patent No. 5470725

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US-09-286-690-8
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GENERAL INFORMATION:
APPLICANT: Li, Xin-Liang
APPLICANT: Liungdahl, Lars G.
APPLICANT: Chen, Huizhong
TITLE OF INVENTION: Lichenase and Coding Sequences
FILE REFERENCE: 55-96
CURRENT APPLICATION NUMBER: US/09/286,690
CURRENT FILING DATE: 1999-04-05
                                                                                                                                                          Sequence 8, Application US/09286690 Patent No. 6103511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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ZIP: 22313-022
COMPUTER READABLE FORM:
REDIUM TYPE: FLOPPy disk
COMPUTER: IBM PC compatible
COMPUTER: PC PCOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 30 TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 836-9300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY & LARDNER
STREET: 1800 Diagonal Road, P.O. Box 299
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TITLE OF INVENTION: A Thermostable (1,3-1,4)-beta-Glucanase
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 16-OCT-1991
ATTORNEY/AGENT INFORMATION:
NAME: Bent, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA: US 07/773,652
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                            190
                                                                                                                                                                                                                                                                                                              143 RKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYKYT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 239 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                    83 EVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEV 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                    23 TNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVD| 82
                                                                                                                                                                                                                                                                                                                                                                                                                              77 TSPSYNKFDCGENRSVQTYGYGLYEVRMKPAKNTGIVSSFFTYTGPTE---GTPWDEID 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Alexandria
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                                                                                                                                                                                                                                                                                                                                                    EFLGKDTTKVQFNYYTNGVGGH----EKVISLGFDASKGFHTYAFDWQPGYIKWYVDG-VL
                                                                                                                                                                                                                                                                            KHT---ATANIPSTPGKIMMNLWNGTGVDDWLGSYNGAN-PLYAEYDWV---KYT
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Olsen, Ole
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Hofemeister, Ju
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13.6%; Score 248; DB 1; Length 239; 35.4%; Pred. No. 5.2e-15;
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US-09-286-690-10
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; TYPE: PRT
; ORGANISM: Bacillus subtilis
US-09-286-690-8
                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 33.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 243
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Best Local Similarity
Matches 63; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Lichenase and Coding Sequences FILE REFERENCE: 55-96 CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05 EARLIER APPLICATION NUMBER: US 60/027,882 EARLIER FILING DATE: 1999-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Li, Xin-Liang
APPLICANT: Ljungdahl, Lars G.
APPLICANT: Chen, Huizhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 12
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EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 24
TYPE: PRT
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                                          143 RKTEGGQVSNLTGTQGLRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYK 193
  194
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     140 QEVRKTEGGQVSNLTGTQGLRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYK 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   134 IDIEFLGKDTTKVQFNYYTNGAG---NHEKIVDLGFDAANAYHTYAFDWQPNSIKWYVDG
                                                                                                                                     83 EVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEV 142
                                                                                                                                                                                   81 TSPSYNKFDCGENRSVQTYGYGLYEVNMKPAKNVGIVSSFFTYTGPT---DGTPWDEIDI 137
                                                                                                                                                                                                                              23 TNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVDI 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              80 VDIEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDG 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           78 ALTSPAYNK-FDCGENRSVQTYGYGLYEVRMKPAKNTGIVSSFFTYTGPT---DGTPWDE 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 ALTTNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVE 79
KHTATTQIPQTPGK--IMMNLWNGAGVDEWLGSYN-GVTPLSRSLHWVRYTK 242
                                                                                          EFLGKDTTKVQFNYYTNGVG---NHEKIVNLGFDAANSYHTYAFDWQPNSIKWYVDGQ-L 193
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                                                                                                                                                                                                                                                                           12.8%; Score 233.5; DB 3 33.7%; Pred. No. 1.1e-13; tive 26; Mismatches 77
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                                                                                                                                                                                                                                                                                                                    DB 3;
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RESULT

Application US/09286690

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SEQ ID NO 9
LENGTH: 279
TYPE: PRT
ORGANISM: Clostridium thermocellum US-09-286-690-9
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US-09-286-690-7
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SEQ ID NO 7
LENGTH: 238
TYPE: PRT
                                                                                 Matches
                                                                                                Query Match
Best Local Similarity
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Best Local Similarity
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EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
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EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/17811
EARLIER FILING DATE: 1997-10-03
                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Li, Xin-Liang
APPLICANT: Ljungdahl, Lars G.
APPLICANT: Chen, Huizhong
TITLE OF INVENTION: Lichenase and Coding Sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              196 VYR----GTRNIPVTPGKIMMNLWPGIGVDEWLGRYD-GRTPLQAEYEYVKYY 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               82
    74
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SLTSPANNK-FDCGEYRSTNNYGYGLYEVSMKPAKNTGIVSSFFTYTGPSH---GTQWDE 129
                         ALTTNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVE 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IEVLGKNPGSFQSNIITGKAGAQKTSEKHAVSPAADQAFHTYGLEWTPNYVRWTVDGQE 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IEFLGKDTTKVQFNWYKNGVGG---NEYLHNLGFDASQDFHTYGFEWRPDYIDFYVDGKK 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ljungdahl, Lars G.
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                                                                                                  12.5%;
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Pred. No. 2.1e-13;
                                                                                                  Score 227.5; DB 3; Pred. No. 3.9e-13;
                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 3;
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TYPE: PRT
; ORGANISM: Clostridium thermocellum
US-09-286-690-11
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                                                                                                                                                                                                                                                                                                            US-08-103-998-4
                                                                                                                                                                                                                                                                                                                              RESULT
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NUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 11
                                                                                                                                                                                                                                                            Sequence 4, Application US/08103998 Patent No. 5470725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Li, Xin-Liang
APPLICANT: Ljungdahl, Lars G.
APPLICANT: Chen, Huizhong
                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Borris:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EARLIER FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: PCT/US97/17811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05 EARLIER APPLICATION NUMBER: US 60/027,882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 55-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Lichenase and Coding Sequences
                                                                                                                                                                        APPLICANT: APPLICANT:
                                                             CORRESPONDENCE ADDRESS:
                                                                                    TITLE OF INVENTION: A Thermostable NUMBER OF SEQUENCES: 4
                                                                                                                             APPLICANT:
                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                         178 SKLPL 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    172 DFHTYGFEWRPDYIDFYVDGKKVYR----GTRNIPVTPGKIMMNLWPGIGVDEWLGRYD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        118 SSFFTYTGPS---DNNPWDEIDIEFLGKDTTKVQFNWYKNGVGG---NEYLHNLGFDASQ 171
                                                                                                                                                                                                                                                                                                                                                                                             227 GRTPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          120 AFHTYGLEWTPNYVRWTVDGQEVRKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDE 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 VAAAAAALTTNVS------AKDFSGAELYTLEEVQ-----YGKFEARMKMAAASGTV 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60 SSMFLYQNGSEIADGRPWVEVDIEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQ 119
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                  ADDRESSEE: FOLEY & LARDNER STREET: 1800 Diagonal Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QEVRKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYKYT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IDIEFLGKDTTKVQFNYYTNGVGGH----EKIINLGFDASTSFHTYAFDWQPGYIKWYVDG 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VSTVLEAFTGDISNGKMILTLDREYGGSYPYKSGEYRTKSFFGYGYYEVRMKAAKNVGIV 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -VLKHT---ATTNIPSTPGKIMMNLWNGTGVDSWLGSYNGAN-PLYAEYDWV---KYT 236
Alexandria
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                                                                                                                             Von Wettstein, Dietrich
                                                                                                                                                      Olsen, Ole
                                                                                                                                                                        Hofemeister, Jurgen
Thomsen, Karl Kristian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                     Thomsen,
                                                                                                                                                                                                                   Borriss, Rainer
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                    P.O.
                      Box 299
                                                                                                        (1,3-1,4)-beta-Glucanase
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; ORGANISM: Orpinomyces sp. PC-2 US-09-286-690-2
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                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/09286690 Patent No. 6103511
                                                                                 SEQ ID NO 2
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                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/286,690 CURRENT FILING DATE: 1999-04-05 EARLIER APPLICATION NUMBER: US 60/027,882 EARLIER FILING DATE: 1999-10-04
                                                                                                                                                                                                                                                                                                            APPLICANT: Li, Xin-Liang
APPLICANT: Ljungdahl, Lars G.
APPLICANT: Chen, Huizhong
                                                                                                                                           EARLIER APPLICATION NUMBER: PCT/US97/17811 EARLIER FILING DATE: 1997-10-03
                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Lichenase and Coding Sequences FILE REFERENCE: 55-96
                                                                                                  SOFTWARE: PatentIn
                                                                                                                            NUMBER OF SEQ ID NOS: 12
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                                          TYPE: PRT
                                                              LENGTH: 245
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LENGTH: 237 amino acids
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/773,652
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       188 KHTATTQIPAAPGK--IMMNLWNGTGVDDWLGSYNGVN-PIYAHYDWMRYRK 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            143 RKTEGGOVSNLTGTQGLRFNLWSSESA-AWVGQFDESKLPLFQFINWVKVYK 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75 TSSAYNKFDCAEYRSTNIYGYGLYEVSMKPAKNTGIVSSFFTYTGP---AHGTQWDEID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23 TNVSAKDESGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVDI
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nes 57; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEV
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899149
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                                                                                                    Ver. 2.0
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; TOPOLOGY: 11; MOLECULE TYPE: US-08-824-707-2
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Matches
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Best Local Similarity
Matches 59; Conserv
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                                                                                                                                                                                                                                                                                                                       TELEFAX: 212-878-9655
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 306 amino acids
                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 212-867-0123
                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION: NAME: Gregg, Valeta A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Savva, Demitris
TITLE OF INVENTION: No. 5919688el enzyme with beta-1,3-glucanase activity
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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 163
                                                                     103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               141 EVRKTEGGQVSNLTGTQG-LRFNLWSSESA-AWVGQFDESKLPLFQFINWV 189
                                  60
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 14
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Gregg, Valeta A. REGISTRATION NUMBER: 35,127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A. ZIP: 10174-6401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81 DIEVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQ 140
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                                                                                                                                                                                                                                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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                                                                                                     N
SAFWMV--GANLPD-TPWPTSGEIDIMENVGNAPHEVHGTVHGPGYSGDNGIMGTYQHPQ 219
                                 SSMFLYQNGSEIADGRPWV---EVDI-EVLGKNP----GSFQSNIITGKAGAQKTSEKHH 111
                                                                   NYTTSRVNSALDGQGNLVITALQESDGSYTSARLTTQGNVQPQFGRIEARIQIPRGQGIW
                                                                                                 NIKKTAVKSALAVAAAAALTTNVSAKDESGAELYTLEEV--QYGKFEARMKMAAASGTV 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AVYTA----YDNIPDTPGKIMMNAWNGIGVDDWLRPFN-GRTNISAYYDWV 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DIEFLGYDTTKVQFNYYTNGQGHH----EHIHYLGFDASQGFHTYGFFWARNSITWYVDGT 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LTIDRDGSGYTCGEYRTKNYYGYGMFQVNMKPIKNPGVVSSFFTYTGPS----DGTKWDEI 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 54;
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                                                                                                                                      Conservative
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xington Avenue, Suite 6400
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27.6%;
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31.6%;
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                                                                                                                                                     Score 159.5; DB Pred. No. 9e-07;
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Pred. No. 9.6e-10;
                                                                                                                                      Mismatches
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                                                                                                                                                                     DB 2;
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                                                                                                                                                                   Length 306;
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LENGTH: 263
TYPE: PRT
                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                         Sequence 11, Application US/09159106 Patent No. 6284509
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Best Local :
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                                                                               APPLICANT: Ferrer, Pau
APPLICANT: Diers, Ivan
APPLICANT: Halkler, Torben
APPLICANT: Hedegaard, Lisbeth
                   TITLE OF INVENTION: An Enzyme With TITLE OF INVENTION: Activity FILE REFERENCE: 4693.204-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EARLIER APPLICATION NUMBER: PCT/DK97/00160
EARLIER FILING DATE: 1997-04-14
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EARLIER APPLICATION NUMBER: 0885/96 EARLIER FILING DATE: 1996-08-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/159,106
CURRENT FILING DATE: 1998-09-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: An Enzyme With TITLE OF INVENTION: Activity FILE REFERENCE: 4693.204-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ferrer, Pau APPLICANT: Diers, Ivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0
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CURRENT APPLICATION NUMBER: US/09/159,106
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QYGRIEARIQIPRGQG-IWPAFWMLGGS--FPGTPWPSGEIDIM-ENVG-FEPHRVHGTV 136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QYGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWVEVDIEVLGKNPGSFQSNIITG-- 99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HGPGYSGGSGITGMYQHPQGWSFADTFHTFAVDWKPGEITWFVDGQQFHRVTRASVG--- 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VGQFDESKLPLFQFINWVKVYKYTPGQGEGGSDF
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                                                                                                                                                                                                                                                                                                                                                                                                                                             GTQGLRFNLWSSESAAWVGQFDESKLPLFQFINWVKVYKYTPGQGEGGS----DFTLDWT 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/09159106
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Pred. No. 1.5e-05;
Pred. No. 1.5e-05;
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Best Local Similarity
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US-09-159-106-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-159-106-13
                                                                                                                                                                                                                         TITLE OF INVENTION: An Enzyme With
TITLE OF INVENTION: Activity
FILE REFERENCE: 4693.204-US
CURRENT APPLICATION NUMBER: US/09/159
CURRENT FILING DATE: 1998-09-23
EARLIER APPLICATION NUMBER: 0427/96
EARLIER FILING DATE: 1996-12-04
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SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 11
LENGTH: 435
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6284509 GENERAL INFORMATION:
                                                                    SEQ ID NO 13
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EARLIER FILING DATE: 1996-12-04
EARLIER FILING DATE: 1996-12-04
EARLIER APPLICATION NUMBER: 0885/96
EARLIER FILING DATE: 1996-08-23
EARLIER APPLICATION NUMBER: PCT/DK97/00160
EARLIER FILING DATE: 1997-04-14
                                                                                                                                                         EARLIER APPLICATION NUMBER: 0885/96
EARLIER FILING DATE: 1996-08-23
EARLIER APPLICATION NUMBER: PCT/DK97/00160
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                                                                                          NUMBER OF SEQ ID NOS: SOFTWARE: FastSEQ for
                                                                                                                                       EARLIER FILING DATE: 1997-04-14
ORGANISM: Oerskovia
                            TYPE: PRT
                                               LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            314 RAANGMCVDVPWADPTDGNPVQIVTCSGNAAQTWTRGS---DGT-VRALGKCLDVRDG 367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      254 QPFFLILNVAVGGQWPGYPDGTTQLPQQMKVDYVRVYDNGSGSSSPGNPGTGLPTGTGAV 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                203 MYQHPQGWSFADTFHTFAVDWKPGEITWFVDGQQFHRVTRASVG-----ANAWVFD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----SAAWVGQFD-ESKLPLFQFINWVKVYKYTPGQGEGGSDFT----
                                               303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09159106
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    xanthineolytica
                                                                                            Windows Version
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Pred. No. 3.9e-05;
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Best Local Similarity
                                                                                                                                                                                    TELEFAX: (212) 286-0854 or 286-0082 TELEX: TWX 710-581-4766 INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIF: 10010
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH 1.44 Mb STORAGE DISKETTE
MEDIUM TYPE: 1BM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPATTING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: you KITY: New York
                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/01
FILING DATE: 11-SEP-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: EN NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
                 ORIGINAL SOURCE:
                                   FRAGMENT TYPE:
                                                 HYPOTHETICAL: I
                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 697-5995
                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           168 SAAWVGQFDESKLPLFQFINWVKVYKYTPGQGEGGS----DFTLDWTDNFDTFDGS 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             203 MYQHPQGWSFADTFHTFAVDWKPGEITWFVDGQQFHRVTRASVG----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             108 EKHHAVSPAADQAFHTYGLEWTPNYVRWTVDGQEVRKTEGGQVSNLTGTQGLRFNLWSSE
                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                      TOPOLOGY:
                                                                                                                   STRANDEDNESS
                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                         NAME: Bogosian, Elizabeth A. REGISTRATION NUMBER: 39,911
                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56 SGTVSSMFLYQNGSEIADGRPWV---EVDI-EVLGKNP----GSFQSNIITGKAGAQKTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88 AELQNYTASRANSALDGQGNLVITARREGDGSYTSARMTTQGKYQPQYGRIEARIQIPRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      34 --- ELYTLE-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28 SALVAALTAAAAALAVTVAATSAAAAPGDLLWSDEFDGAAGSAPNPAVWNHETGAHGWGN
                                                                                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6 TAVKSALAVAAAAALT---TNVSA-----KDFSGA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ANAWV--FDQ---PFFLILN-VAVGGQWPGYPDGTTQLPQQMKVDYVRVYDNGSGS 290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QG-IWPAFWMLGGS--FPGTPWPSSGEIDIMENVGFEPHRVHGTVHGPGYSGGSGI--TG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10016
                                                                                                                                   amino acid
                                                                                                                                                      276 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90 Park Avenue
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gub, Rhodothermus marinus
                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amster, Rothstein & Ebenstein
                                                                                   peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Goldstein, Alex Zhu and Lin Leng DN: ENDO-BETA-GALACTOSIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.8%; Score 141.5;
23.0%; Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/712,072C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34;
                                                                                                                                                                                                                                                                           63475/97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -----EVQYGKFEARMKMAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               167
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RESULT 15
US-08-712-072C-3
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                                                                                                                                                                                                                                                                                                                          Matches
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                                                                  269 VRVYRW 274
                                                                                                189 VKVYKY 194
                                                                                                                                 209 DDSLYYRFPNERLTDPEADWRHWPFDQPFHLIMNIAVGGAWGGQQGVDPEAFPAQLVVDY 268
                                                                                                                                                                 138
                                                                                                                                                                                               152 MEHVGFNPDVVHGTVHTKAYNHLLGTQRGGSIR---VPTARTDFHVYAIEWTPEEIRWFV
                                                                                                                                                                                                                           83 -EVLGKNP----GSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLEWTPNYVRWTV 137
                                                                                                                                                                                                                                                            93 REYTSARLVTRGKASWTYGRFEIRARLPSGRGTWPAIWMLPDRQTYGSAY-WPDNGEIDI 151
                                                                                                                                                                                                                                                                                          28 KDFSGAELYTLEEVQ--YGKFEARMKMAAASGTVSSMFLYQNGSEIADGRPWV---EVDI 82
                                                                                                                                                                                                                                                                                                                        Local Similarity
nes 44; Conserv
                                                                                                                                                             DGQEVRKTEGGQVSNLTG--
                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                         7.3%;
                                                                                                                                                               ----TQGLRFNLWSSESAAWVGQ--FDESKLPLFQFINW 188
                                                                                                                                                                                                                                                                                                                        34; Mismatches
                                                                                                                                                                                                                                                                                                                                            Pred. No.
                                                                                                                                                                                                                                                                                                                                                         Score 133.5;
                                                                                                                                                                                                                                                                                                                                         0.00018;
                                                                                                                                                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                      Length 276;
                                                                                                                                                                                                                                                                                                                          23;
                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                               208
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Sequence 3, Application US/08712072C Patent No. 5925541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Jack Gold
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                           TELEX: TWX 710-581-4766 INFORMATION FOR SEQ ID NO:
FRAGMENT TYPE:
ORIGINAL SOURCE:
ORGANISM: e131
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH 1.44 N
COMPUTER: IBM PC COMPATIBLE
                                                 HYPOTHETICAL: | ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION NAME: Bogosian, Elizabet
                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS:
                                                                                                 TOPOLOGY:
                                                                                                                                                                                                            TELEFAX:
                                                                                                                                                                                                                          TELEPHONE:
                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 63475/97
                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10016
                                                                                                                                                                                                                                                                                      Bogosian, Elizabeth A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ΝY
                                                                                                                                             321 amino acids
                                                                                                                                                                                                             (212)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jack Goldstein,
   e13b,
                                                                                                 linear
                                                                                                                                                                                                                         (212)
                                                                                                                                                                                                                                                                                                                                                                                     11-SEP-1996
                                                                                                                                                                                                            286-0854 or 286-0082
 Bacillus circulars
                                                                                                                                                                                                                            697-5995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INCH 1.44 Mb STORAGE DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                    MS-DOS
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ENDO-BETA-GALACTOSIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rothstein & Ebenstein
                                                                                                                                                                                                                                                                                                                                                                                                      US/08/712,072C
                                                                                                                                                                                                                                                                     39,91
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Sear Job	DЪ	Оу	Qу	ОУ	Qu <b>x</b> a
ch co time:	170 290	128 244	72 184	20 124	Query Match Best Local   Matches 4
Search completed: January 15, 2002, 15:53:08 Job time: 288 sec	170 AWVGQFDESKLPLFQFINWVKVYKY 194	128 WTPNYVRWTVDGQEVRKTEGGQVSNLTGTQGLRFNLWSSESA 169.   : :      :	72 ADGRPWVEVDI-EVLGKNPGSFQSNIITGKAGAQKTSEKHHAVSPAADQAFHTYGLE 127  : :    :    :    :    :	20 ALTTNVSAKDFSGAELYTLEEVQYGKFEARMKMAAASGTVSSMFLYQNGSEI 71	Query Match 6.5%; Score 119; 'DB 2; Length 321; Best Local Similarity 20.4%; Pred. No. 0.0048; Matches 43; Conservative 34; Mismatches 84; Indels 50; Gaps 7;

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